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<210> 7
<211> 65
<212> DNA
<213> Homo sapiens

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<400> 7
agtcctcgtg ctcttagccc tcttcacaa caggaaacca atatgattag tttctttcat 60
aggct 65

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<210> 8
<211> 656
<212> DNA
<213> Homo sapiens

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<400> 8
gtgcagcctc agggcgccgc ctctcgacct tcccgcctcc acctcccacc gcccgccctc 60
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ggcccatggg ggagccgggt cgccactccc ggaccgccc cctcgagggg ggtggagctg 180
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cccaaggata ggaaggttca ggcaaccggc tggccgtgtc ttgggggctt cattgctggg 540
caaaggcagt gcagcagacg gagacaaact ttcttccctg gccgtggcca gagggcagaa 600
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<210> 9
 <211> 177
 <212> DNA
 <213> Homo sapiens

<400> 9
 cggggcacggg tcggccgcaa tccagcctgg gcggagccgg agttgcgagc cgctgcctag 60
 aggcgcgagg gctcacagct atggggtgga gggcccggag agctcggggg accccgttgc 120
 tgtgctgctg actactgctg ctgctctggc cagtgccagg cgcgggggtg cttcaag 177

<210> 10
 <211> 80
 <212> DNA
 <213> Homo sapiens

<400> 10
 gacatatccc tgggcagcca gtcaccccgc actgggtcct ggatggacaa ccttggcgca 60
 ccgtcagcct ggaggagccg 80

<210> 11
 <211> 77
 <212> DNA
 <213> Homo sapiens

<400> 11
 gtctcgaagc cagacatggg gctgggtggc ctggaggctg aaggccagga gctcctgctt 60
 gagctggaga agaacca 77

<210> 12
 <211> 79
 <212> DNA
 <213> Homo sapiens

<400> 12
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 gctggccccc aaccacacg 79

<210> 13
 <211> 119
 <212> DNA
 <213> Homo sapiens

<400> 13
 cagggtgctg gcccaggat acatagaaac ccaactacggc ccagatgggc agccagtggg 60
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<210> 14
 <211> 77
 <212> DNA
 <213> Homo sapiens

<400> 14

gatcattgcc actaccaagg gogagtaagg ggcttccccg actcctgggt agtcctctgc 60
acctgctctg ggatgag 77

<210> 15
<211> 190
<212> DNA
<213> Homo sapiens

<400> 15
tggcctgac accctcagca ggaatgccag ctattatctg cgtccctggc caccctgggg 60
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aacctgtggc cacagggatc ctgggaacaa agcgggcatg accagccttc ctggtggtcc 180
ccagagcagg 190

<210> 16
<211> 66
<212> DNA
<213> Homo sapiens

<400> 16
ggcagggcag aagcgcgcag gacccggaag tacctggaac tgtacattgt ggcagaccac 60
accctg 66

<210> 17
<211> 72
<212> DNA
<213> Homo sapiens

<400> 17
ttcttgactc ggcaccgaaa cttgaaccac accaaacagc gtctcctgga agtcgccaac 60
tacgtggacc ag 72

<210> 18
<211> 167
<212> DNA
<213> Homo sapiens

<400> 18
cttctcagga ctctggacat tcaggtggcg ctgaccggcc tggaggtgtg gaccgagcgg 60
gaccgcagcc gcgtcacgca ggacgccaac gccacgctct gggccttcct gcagtgggcg 120
cgggggctgt gggcgacgag gccccacgac tccgcgcagc tgctcac 167

<210> 19
<211> 85
<212> DNA
<213> Homo sapiens

<400> 19
gggcccgcgc ttccagggcg ccacagtggg cctggcgccc gtcgagggca tgtgccgcgc 60
cgagagctcg ggaggcgtga gcacg 85

<210> 20

<211> 143
 <212> DNA
 <213> Homo sapiens

<400> 20
 gaccactcgg agctccccat cggcgccgca gccaccatgg cccatgagat cggccacagc 60
 ctccggcctca gccacgaccc cgacggctgc tgcgtggagg ctgcggccga gtccggagggc 120
 tgcgtcatgg ctgcggccac cgg 143

<210> 21
 <211> 178
 <212> DNA
 <213> Homo sapiens

<400> 21
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 caaggggggg ggcgcttgcc tetccaatgc cccggacccc ggactcccg tgcgcggcgc 120
 gctctgcggg aacggcttcg tgggaagcgg cgaggagtgt gactgcggcc ctggccacg 178

<210> 22
 <211> 90
 <212> DNA
 <213> Homo sapiens

<400> 22
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 gcccacgggg actgctgcgt gcgctgcctg 90

<210> 23
 <211> 196
 <212> DNA
 <213> Homo sapiens

<400> 23
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 cagctctggg ggcctg 196

<210> 24
 <211> 107
 <212> DNA
 <213> Homo sapiens

<400> 24
 gctccacccc agctcccgag gcctgtttcc aggtgggtgaa ctctgcggga gatgetcatg 60
 gaaactgcgg ccaggacacg gaggggccact tcctgcctcg tgcaggg 107

<210> 25
 <211> 199
 <212> DNA
 <213> Homo sapiens

<400> 25
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 catggtgcca gtggactcta ccgttcacct agatggccag gaagtgcatt gtcggggagc 120
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 ccagtggtga cctagaatg 199

<210> 26
 <211> 109
 <212> DNA
 <213> Homo sapiens

<400> 26
 gtttgcataa gcaaccataa ctgccactgt gctccaggct gggctccacc cttctgtgac 60
 aagccaggct ttggtggcag catggacagt ggccctgtgc aggtgaaa 109

<210> 27
 <211> 148
 <212> DNA
 <213> Homo sapiens

<400> 27
 accatgacac cttctctgtg gccatgctcc tcagcgtcct gctgcctctg ctcccagggg 60
 ccggcctggc ctggtgttgc taccgactcc caggagccca tctgcagcga tgcagctggg 120
 gctgcagaag ggaccctgag tgcagtgg 148

<210> 28
 <211> 92
 <212> DNA
 <213> Homo sapiens

<400> 28
 ccccaaagat ggcccacaca gggaccaccc cctggggcggc gttcacccca tggagtggg 60
 cccacagcc actggacagc cctggccctt gg 92

<210> 29
 <211> 72
 <212> DNA
 <213> Homo sapiens

<400> 29
 accctgagaa ctctcatgag cccagcagcc accctgagaa gcctctgcca gcagtctcgc 60
 ctgaccccca ag 72

<210> 30
 <211> 1031
 <212> DNA
 <213> Homo sapiens

<400> 30
 cagatcaagt ccagatgcca agatcctgccc tctgggtgaga ggtagctcct aaaatgaaca 60
 gatttaaaga caggtggcca ctgacagcca ctccaggaaac ttgaactgca ggggcagagc 120
 cagtgaatca ccggacctcc agcacctgca ggcagcttgg aagtttcttc cccagtgga 180
 gcttcgaccc accactcca ggaaccacga gccacattag aagttcctga gggctggaga 240

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acactgctgg gcacactctc cagctcaata aaccatcagt cccagaagca aaggtcacac 300
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aagacataaa a 1031

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<210> 31
<211> 78
<212> DNA
<213> Homo sapiens

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<400> 31
gtgtgccaga gcaggcgtg caggaagaat gccttcagg agcttcagcg ctgcctgact 60
gcctgccaca gccacggg 78

```

```

<210> 32
<211> 6
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence: polyhistidine tag

```

```

<400> 32
His His His His His His
1 5

```

```

<210> 33
<211> 8
<212> PRT
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence: FLAG epitope tag

```

```

<400> 33
Asp Tyr Lys Asp Asp Asp Asp Lys
1 5

```

```

<210> 34
<211> 22
<212> DNA
<213> Artificial Sequence

```

```

<220>

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<223> Description of Artificial Sequence: Primer

<400> 34

aactcttgaa atgagaagcg tg

22

<210> 35

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 35

aatatcatgc accatgaccc ac

22

<210> 36

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 36

tggagtaaagt attgtaaact at

22

<210> 37

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 37

ggagettatc ctggattatc ta

22

<210> 38

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 38

agagccacac atccatgtcc tg

22

<210> 39

<211> 22

<212> DNA

<213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

 <400> 39
 aagccactct gtgaattgcc at 22

 <210> 40
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 40
 gagtagtcgt agtaccagat gg 22

 <210> 41
 <211> 22
 <212> DNA
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 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 41
 gtctggcaat ggagcatgaa aa 22

 <210> 42
 <211> 22
 <212> DNA
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 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 42
 attagagcac atgaaggaaa gg 22

 <210> 43
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 43
 aactgcttt gggggacagg ct 22

 <210> 44
 <211> 22

<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer

<400> 44
cacgacgccacagaccagc tc 22

<210> 45
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 45
aaccaccacg gattcacgct tc 22

<210> 46
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 46
ataaccagat ggctgtgggt ca 22

<210> 47
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 47
atccccgcaa tgaaatagtt ta 22

<210> 48
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 48
gttgagagcc cacttagata at 22

<210> 49
 <211> 22
 <212> DNA
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<220>
 <223> Description of Artificial Sequence: Primer

<400> 49
 gcattggggg aagccaggac at 22

<210> 50
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 50
 gccactagga ggcaatggca at 22

<210> 51
 <211> 22
 <212> DNA
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<220>
 <223> Description of Artificial Sequence: Primer

<400> 51
 cgacggcatc acggccatct gg 22

<210> 52
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 52
 tccaggctca ttcattttca tg 22

<210> 53
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 53
 tgacatcaac ttctcctttc ct 22

<210> 54
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 54
 agttgcagag acctagcctg tc 22

<210> 55
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 55
 tctgggagag gacggagctg gc 22

<210> 56
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 56
 tgtaggacta tattgctc 18

<210> 57
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 57
 cgacatttag gtgacact 18

<210> 58
 <211> 15
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: BstXI-linker
 adapter

<400> 58
gttttcacca cgggg

15

<210> 59
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: BstXI-linker
adapter

<400> 59
gtggagaaga c

11

<210> 60
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
peptide

<400> 60
Asp Pro Gln Ala Asp Gln Val Gln Met
1 5

<210> 61
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
peptide

<400> 61
Asp Pro Gln Asp Gln Val Gln Met
1 5

<210> 62
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<221> MOD_RES
<222> (1)..(11)
<223> "Xaa" represents a variable amino acid

<220>
<223> Description of Artificial Sequence: Zn-binding

consensus sequence

<400> 62

His Glu Xaa Xaa His Xaa Xaa Gly Xaa Xaa His

1 5 10

<210> 63

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 63

ctgcctagag gccgagga

18

<210> 64

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 64

caggagacca cggaagatcg

20

<210> 65

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 65

ttgcctgaac cttcctatcc

20

<210> 66

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 66

ccctgtgtt cctcaggtc

19

<210> 67

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 67

gctccacact ctttcttgcc

20

<210> 68

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 68

aggcaggagg aagctgaat

19

<210> 69

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 69

cctaccacac cctcccttt

20

<210> 70

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 70

cctacccttc tgcacccta

19

<210> 71

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 71

aacttccttc tgggagctgg

20

<210> 72

<211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 72
 cacaccctgg tgaggagaga 20

<210> 73
 <211> 16
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 73
 ccacgaagga ccaccg 16

<210> 74
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 74
 ctcacgtggg tgcctctg 18

<210> 75
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 75
 ctctacggcc gcagtgac 18

<210> 76
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 76
 gtccctccat gcccaatg 18

<210> 77
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 77
cagggttaagt cggctcgc 18

<210> 78
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 78
ctctctctgc cttcccccac 19

<210> 79
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 79
tctactgtgg ggaagatggg 20

<210> 80
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 80
cccctctact tcctcccca 19

<210> 81
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 81

gaccttgggg ttcctaatacc

20

<210> 82

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 82

gtgcacctgc tcaggactc

19

<210> 83

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 83

cctggactct tatcacgttg c

21

<210> 84

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 84

ttaccctcca ccatttctcc

20

<210> 85

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 85

gtggagaggg aagggagaag

20

<210> 86

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 86
 ccccatgggt tgaatttaca 20

<210> 87
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 <220>
 <223> Description of Artificial Sequence: Primer

<400> 87
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<210> 88
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 <220>
 <223> Description of Artificial Sequence: Primer

<400> 88
 accacgccta tagccaacat 20

<210> 89
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Primer

<400> 89
 aggtgtagca ctgggattgg 20

<210> 90
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Primer

<400> 90
 cccaggacc actagcttct 20

<210> 91
 <211> 20
 <212> DNA
 <213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Primer

<400> 91
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His Gly Asp Cys Cys Val Arg Cys Leu Lys Pro Ala Gly Ala Leu
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Cys Arg Gln Ala Met Gly Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly
 35 40 45

Thr Ser Ser His Cys Pro Pro
 50 55

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 20 25 30

Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
 35 40 45

Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
 50 55 60

Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
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Leu Glu Lys Asn His Arg
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Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
35 40 45

Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
50 55 60

Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
65 70 75 80

Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
85 90 95

Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Val
100 105 110

Arg Cys Phe His Gly Leu Trp Asp Ala Pro Pro Glu Asp His Cys His
115 120 125

Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val Val Leu Cys
130 135 140

Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg Asn Ala Ser
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Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp Phe Ser Thr
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His Glu

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Phe Ser Thr His Glu
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35 40 45
Met Ser Gly Leu Ile Thr Leu Ser Arg Asn Ala Ser Tyr Tyr Leu Arg
50 55 60
Pro Trp Pro Pro Arg Gly Ser Lys Asp Phe Ser Thr His Glu Ile Phe
65 70 75 80
Arg Met Glu Gln Leu Leu Thr Trp Lys Gly Thr Cys Gly His Arg Asp
85 90 95
Pro Gly Asn Lys Ala Gly Met Thr Ser Leu Pro Gly Gly Pro Gln Ser
100 105 110
Arg Gly Arg Arg Lys Ala Arg Arg Thr Arg Lys Tyr Leu Glu Tyr
115 120 125
Ile Val Ala Asp His Thr Leu Phe Leu Thr Arg His Arg Asn Leu Asn
130 135 140
His Thr Lys Gln Arg Leu Leu Glu Val Ala Asn Tyr Val Asp Gln Leu
145 150 155 160
Leu Arg Thr Leu Asp Ile Gln Val
165

<210> 343
<211> 167
<212> PRT
<213> Homo sapiens

<400> 343
Ser Gly Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys
1 5 10 15
Gln Gln Leu Trp Gly Pro Gly Ser His Pro Ala Pro Glu Ala Cys Phe
20 25 30
Gln Val Val Asn Ser Ala Gly Asp Ala His Gly Asn Cys Gly Gln Asp
35 40 45

Ser Glu Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly
 50 55 60
 Lys Leu Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met
 65 70 75 80
 Val Pro Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys
 85 90 95
 Arg Gly Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu
 100 105 110
 Gly Leu Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Asn
 115 120 125
 Ser Asn His Asn Cys His Cys Ala Pro Gly Trp Ala Pro Pro Phe Cys
 130 135 140
 Asp Lys Pro Gly Phe Gly Gly Ser Met Asp Ser Gly Pro Val Gln Ala
 145 150 155 160
 Glu Asn His Asp Thr Phe Leu
 165
 <210> 344
 <211> 193
 <212> PRT
 <213> Homo sapiens
 <400> 344
 Ser Gly Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys
 1 5 10 15
 Gln Gln Leu Trp Gly Pro Gly Ser His Pro Ala Pro Glu Ala Cys Phe
 20 25 30
 Gln Val Val Asn Ser Ala Gly Asp Ala His Gly Asn Cys Gly Gln Asp
 35 40 45
 Ser Glu Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly
 50 55 60
 Lys Leu Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met
 65 70 75 80
 Val Pro Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys
 85 90 95
 Arg Gly Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu
 100 105 110
 Gly Leu Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln
 115 120 125
 Ser Arg Arg Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu
 130 135 140

Thr Ala Cys His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His
145 150 155 160

Cys Ala Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly
165 170 175

Gly Ser Met Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe
180 185 190

Leu

<210> 345

<211> 126

<212> PRT

<213> Homo sapiens

<400> 345

Ser Gly Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys
1 5 10 15

Gln Gln Leu Trp Gly Pro Asp Gly Gln Glu Val Thr Cys Arg Gly Ala
20 25 30

Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val
35 40 45

Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg
50 55 60

Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys
65 70 75 80

His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro
85 90 95

Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met
100 105 110

Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu
115 120 125

<210> 346

<211> 93

<212> PRT

<213> Homo sapiens

<400> 346

Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser
1 5 10 15

Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro
20 25 30

His Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro
35 40 45

Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu
50 55 60

Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro
65 70 75 80

Gln Ala Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp
85 90

<210> 347

<211> 236

<212> PRT

<213> Homo sapiens

<400> 347

Ser Gly Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys
1 5 10 15

Gln Gln Leu Trp Gly Pro Asp Gly Gln Glu Val Thr Cys Arg Gly Ala
20 25 30

Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val
35 40 45

Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg
50 55 60

Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys
65 70 75 80

His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro
85 90 95

Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met
100 105 110

Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu Ala
115 120 125

Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu Ala
130 135 140

Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp
145 150 155 160

Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His
165 170 175

Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro Thr
180 185 190

Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro
195 200 205

Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln
210 215 220

400> 348	Ser	Gly	Tyr	Cys	Trp	Asp	Gly	Ala	Cys	Pro	Thr	Leu	Glu	Gln	Gln	Cys
1					5					10					15	
Gln	Gln	Leu	Trp	Gly	Pro	Gly	Ser	His	Pro	Ala	Pro	Glu	Ala	Cys	Phe	
			20					25						30		
Gln	Val	Val	Asn	Ser	Ala	Gly	Asp	Ala	His	Gly	Asn	Cys	Gly	Gln	Asp	
		35					40					45				
Ser	Glu	Gly	His	Phe	Leu	Pro	Cys	Ala	Gly	Arg	Asp	Ala	Leu	Cys	Gly	
	50					55					60					
Lys	Leu	Gln	Cys	Gln	Gly	Gly	Lys	Pro	Ser	Leu	Leu	Ala	Pro	His	Met	
	65				70					75					80	
Val	Pro	Val	Asp	Ser	Thr	Val	His	Leu	Asp	Gly	Gln	Glu	Val	Thr	Cys	
			85					90						95		
Arg	Gly	Ala	Leu	Ala	Leu	Pro	Ser	Ala	Gln	Leu	Asp	Leu	Leu	Gly	Leu	
		100						105					110			
Gly	Leu	Val	Glu	Pro	Gly	Thr	Gln	Cys	Gly	Pro	Arg	Met	Val	Cys	Gln	
	115						120					125				
Ser	Arg	Arg	Cys	Arg	Lys	Asn	Ala	Phe	Gln	Glu	Leu	Gln	Arg	Cys	Leu	
	130					135					140					
Thr	Ala	Cys	His	Ser	His	Gly	Val	Cys	Asn	Ser	Asn	His	Asn	Cys	His	
145					150					155					160	
Cys	Ala	Pro	Gly	Ala	Pro	Pro	Phe	Cys	Asp	Lys	Pro	Gly	Phe	Gly		
			165					170					175			
Gly	Ser	Met	Asp	Ser	Gly	Pro	Val	Gln	Ala	Glu	Asn	His	Asp	Thr	Phe	
			180					185					190			
Leu	Leu	Ala	Met	Leu	Leu	Ser	Val	Leu	Leu	Pro	Leu	Leu	Pro	Gly	Ala	
	195						200					205				
Gly	Leu	Ala	Trp	Cys	Cys	Tyr	Arg	Leu	Pro	Gly	Ala	His	Leu	Gln	Arg	
	210					215					220					
Cys	Ser	Trp	Gly	Cys	Arg	Arg	Asp	Pro	Ala	Cys	Ser	Gly	Pro	Lys	Asp	
	225				230					235					240	
Gly	Pro	His	Arg	Asp	His	Pro	Leu	Gly	Val	His	Pro	Met	Glu	Leu		
			245					250					255			

Gly Pro Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser
260 265 270

His Glu Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro
275 280 285

Asp Pro Gln Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp
290 295 300

<210> 349

<211> 235

<212> PRT

<213> Homo sapiens

<400> 349

Ser Gly Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys
1 5 10 15

Gln Gln Leu Trp Gly Pro Asp Gly Gln Glu Val Thr Cys Arg Gly Ala
20 25 30

Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val
35 40 45

Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg
50 55 60

Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys
65 70 75 80

His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro
85 90 95

Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met
100 105 110

Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu Ala
115 120 125

Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu Ala
130 135 140

Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp
145 150 155 160

Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His
165 170 175

Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro Thr
180 185 190

Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro
195 200 205

Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln
210 215 220

Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp
 225 230 235

<210> 350
 <211> 339
 <212> DNA
 <213> Homo sapiens

<400> 350
 cgggacacggg tcggccgcaa tccagcctgg gcggagccgg agttgcgagc cgctgcctag 60
 aggcgcgagga gctcacagct atgggctgga ggcccgagg agctcggggg accccggttg 120
 tgctgctgct actactgctg ctgctctggc cagtgccagg cgccgggggtg cttcaaggac 180
 atatccctgg gcagccagtc accccgcact gggctcctgga tggacaacct tggcgccacc 240
 tcagcctgga ggagccggtc tcgaagccag acatgggggt ggtggccctg gaggctgaag 300
 gccaggagct cctgcttgag ctggagaaga accacaggc 339

<210> 351
 <211> 225
 <212> DNA
 <213> Homo sapiens

<400> 351
 cgggacacggg tcggccgcaa tccagcctgg gcggagccgg agttgcgagc cgctgcctag 60
 aggcgcgagga gctcacagct atgggctgga ggcccgagg agctcggggg accccggttg 120
 tgctgctgct actactgctg ctgctctggc cagtgccagg cgccgggggtg cttcaaggac 180
 atatccctgg gcagccagtc accccgcact gggctcctgga tggac 225

<210> 352
 <211> 562
 <212> DNA
 <213> Homo sapiens

<400> 352
 gcctagaggc cgaggagctc acagctatgg gctggaggcc ccggagagct cgggggacct 60
 cgttgctgct gctgctaacta ctgctgctgc tctggccagt gccaggcgcc ggggtgcttc 120
 aaggacatat ccttggggcag ccagtcaccc cgcactgggt cctggatgga caacctggc 180
 gcaccgtcag cctggaggag ccggctctga agccagacat ggggctgggt gccctggagg 240
 ctgaaggcca ggagctcctg cttgagctgg agaagaacca caggctgctg gccccaggat 300
 acatagaaac ccaatcagcc ccagatgggc agccagtggt gctggccccc aaccacacgg 360
 tggatgctct ccaatgggctc tgggatgcac cgccagagga tcattgccac taccaggggc 420
 gagtaagggg cttcccccac tcctgggtag tcctctgcac ctgctctggg atgagtggcc 480
 tgatcacctc cagcaggaat gccagctatt atctgcgtcc ctggccacct cggggctcca 540
 aggaactctc aaccacagag at 562

<210> 353
 <211> 362
 <212> DNA
 <213> Homo sapiens

<400> 353
 gaggccgagg agctcacagc tatgggctgg agggcccgga gagctcgggg gaccccggtg 60
 ctgctgctgc tactactgct gctgctctgg ccagtgccag gcgccgggtg gcttcaagga 120
 cataccctg ggcagccagt caaccgcac tgggtcctgg atggacaacc ctggcgccacc 180

```

gtcagcctgg aggagccggt ctccaagcca gacatggggc tgggtggccct ggaggctgaa 240
ggccaggagag tctgtcttga gctggagaag aaccatggcc tgatcacctt cagcaggaaat 300
gccagctatt atctgcgtcc ctggccaccc cgggggtcca aggacttctc aacccacgag 360
at

```

<210> 354

<211> 518

<212> DNA

<213> Homo sapiens

<400> 354

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gaggccgagg agctcacagc tatgggctgg agggcccgga gagctcgggg gaccccggtg 60
ctgctgctgc tactactgct gctgctctgg ccagtgccag gcgcgggggt gcttcaaggga 120
catatccctg ggcagccagt caccocgcac tgggtcctgg atggacaacc ctggcgacc 180
gtcagcctgg aggagccggt ctccaagcca gacatggggc tgggtggccct ggaggctgaa 240
ggccaggagag tctgtcttga gctggagaag aaccacaggg tgctggccctc aggatacata 300
gaaacccact acggccccaga tgggcagcca gtggtgctgg cccccaacca caccgatcat 360
tgccaactacc aagggcgagg aaggggcttc cccgactcct gggtagtctt ctgcaactgc 420
tctgggatga gtggcctgat caccctcagc aggaatgccg gctattatct gcgtccctgg 480
ccaccccggg gctccaagga cttctcaacc cagcagat

```

<210> 355

<211> 506

<212> DNA

<213> Homo sapiens

<400> 355

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ctggcccccag gatacataga aaccactac gggccagatg ggcagccagt ggtgctggcc 60
cccaaccaca cggatcatgt ccactaccaa gggcgagtaa ggggcttccc cgactcctgg 120
gtagtctctc gcacctgctc tgggatgagt ggcctgatca cctcagcag gaatgcacag 180
tattatctgc gtccctggcc accccggggc tccaaggact tctcaacca cgagatcttt 240
cggatggagg agctgctcac ctggaaggga acctgtggcc acagggatcc tgggaacaaa 300
gcgggcatga ccagccttcc tgggtggtccc cagagcaggg gcagggcgaag agcgcgagg 360
accgggaagt acctggaact gtacattgtg gcagaccaca cctgtgtctt gactcgccac 420
cgaaaacttga accacaccaa acagcgtctc ctggaagtgc ccaactacgt ggaccagctt 480
ctcaggactc tggacattca ggtggc

```

<210> 356

<211> 503

<212> DNA

<213> Homo sapiens

<400> 356

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cagtggtctac tggctgggatg gcgcattgtc cagcgtggag cagcagtgcc agcagctctg 60
ggggcctggc tcccaccagc ctcccagggc ctgtttccag gtggtgaact ctgcgggaga 120
tgctcatgga aactgcggcc aggacagcga gggccacttc ctgcccctgtg caggggaggga 180
tgccctgtgt ggggaagctgc agtgccaggg tggaaagccc agcctgtctg caccgcacat 240
gggtccagtg gactctacgc ttacactaga tggccaggaa gtgacttgtc ggggagacctt 300
ggcactcccc agtgccacag tggacctgct tggcctgggc ctggttagagc caggacacca 360
gtgtggacct agaattggtt gcaatagcaa ccataactgc cactgtgctc caggctgggc 420
tccacccttc tgtgacaagc caggctttgg tggcagcagt gacagtgggc ctgtgcaggc 480
tgaaaaccat gacaccttcc tgc

```

<210> 357

<211> 581
 <212> DNA
 <213> Homo sapiens

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<400> 357
cagtggctac  tgctgggatg  gcgcattgtcc  cacgctggag  cagcagtgcc  agcagctctg  60
ggggcctggc  tcccacccag  ctcccaggcc  ctgtttccag  gtggtgaaat  ctgcgggaga  120
tgctcatgga  aactcgggcc  aggacagcga  gggcacttc  ctgcctgtg  cagggaggga  180
tgccctgtgt  ggggaagctg  agtgccaggg  tggaaaagcc  agcctgctg  caccgcacat  240
ggtgccatgt  gactctaccg  ttacactaga  tggccaggaa  gtgacttgct  ggggagcctt  300
ggcaactccc  agtgcccagc  tggacctgct  tggcctgggc  ctggtagagc  caggcaccca  360
gtgtggacct  agaatggtgt  gccacagcag  gcgctgcagg  aagaatgcct  tccaggagct  420
tcagcgctgc  ctgactgcct  gccacagcca  cgggggttgc  aatagcaacc  ataactgcca  480
ctgtgtccca  ggctgggctc  caccctctgt  tgacaagcca  ggctttggtg  gcagcagtga  540
cagtggccct  gtgcaggctg  aaaacccatga  caccctctgt  c
```

<210> 358
 <211> 380
 <212> DNA
 <213> Homo sapiens

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<400> 358
cagtggctac  tgctgggatg  gcgcattgtcc  cacgctggag  cagcagtgcc  agcagctctg  60
ggggcctgat  gggcaggaa  tgacttctgt  gggagccttg  gcaactccca  gtgccagct  120
ggacctgctt  ggctcgggcc  tggtagagcc  aggacccag  tgtggacct  gaattggtg  180
ccagagcagg  cgctgcagg  agaattgcct  ccaggagctt  cagcgctgc  tgactgcctg  240
ccacagccac  gggggttgca  atagcaacca  taactgccac  tgtgctccag  gctgggctcc  300
accctctctg  gacaagccag  gctttggttg  cagcattggc  agtggccctg  tgcaggctga  360
aaacctgatc  accctcctgc
```

<210> 359
 <211> 324
 <212> DNA
 <213> Homo sapiens

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<400> 359
ggcctgggtg  tgctaccgac  tcccaggagc  ccactctcag  cgatgcagct  ggggctgcag  60
aagggaacct  gcgtgcagtg  gccccaaga  tggcccaac  agggaccaac  cctgggggg  120
cgttcacccc  atggagtgg  gccccacagc  cactggacag  ccttgcccc  tggacctga  180
gaactctcat  gagcccagca  gccaccccta  gaagcctctg  ccagcactct  cgctgaccc  240
ccaagcagat  caagtccaga  tgcacaagtc  ctgcctcttg  tgagaggtag  ctccataaat  300
gaacagattt  aaagacaggt  gcc
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<210> 360
 <211> 753
 <212> DNA
 <213> Homo sapiens

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<400> 360
cagtggctac  tgctgggatg  gcgcattgtcc  cacgctggag  cagcagtgcc  agcagctctg  60
ggggcctgat  gggcaggaa  tgacttctgt  gggagccttg  gcaactccca  gtgccagct  120
ggacctgctt  ggctcgggcc  tggtagagcc  aggacccag  tgtggacct  gaattggtg  180
ccagagcagg  cgctgcagg  agaattgcct  ccaggagctt  cagcgctgc  tgactgcctg  240
ccacagccac  gggggttgca  atagcaacca  taactgccac  tgtgctccag  gctgggctcc  300
accctctctg  gacaagccag  gctttggttg  cagcattggc  agtggccctg  tgcaggctga  360
```

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aaaccatgac accctcctgc tggccatgct cctcagcgtc ctgctgcctc tgctcccagg 420
ggccggcctg gcttgggtgt gctaccgact cccaggagcc catctgcagc gatgcagctg 480
gggctgcaga agggaccctg cgtgcagttg ccccaaagat ggcccacaca gggaccaccc 540
cctggggcgg gttcacccca tggagttggg cccacagccc actggagccc cctggccctc 600
ggaccctcag aactctcatg agcccagcag ccaccctgag aagcctctgc cagcagcttc 660
gctgcacccc caagcacatc aagtcagat gccaatatcc tgctctgggt gagaggtagc 720
tcctaaaaat aacagattta aagacaggtg gcc 753

```

<210> 361

<211> 1154

<212> DNA

<213> Homo sapiens

<400> 361

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cagtggctac tgetgggatg gcgcatgtcc cacgctggag cagcagtgcc agcagctctg 60
ggggcctggc tcccaccagc ctcccaggcc ctgtttccag gtgggaact ctgcgggaga 120
tgctcatgga aactcggccc aggcacagcga gggccacttc ctgccctgtg caggagaggga 180
tgccctgtgt gggaaagctg agtgccaggg tggaaaagccc agcctgctgc caccgcacat 240
ggtgccagtg gactctaccg ttcacctaga tggccaggaa gtgactgtgc ggggagcctt 300
ggcactcccc agtgcccagc tggacctgct tggcctgggc ctgttagagc caggacacca 360
gtgtggacct agaattggtg gccagagcag gcgctgcagg aagaatgcct tcaggagact 420
tcagcgtctg ctgactgcct gccacagcca cggggtttgc aatagaaccc ataatgcaca 480
ctgtgctcca ggtgggcttc cacccttctg tgacaagcca ggctttgggt gcagcatgga 540
cagtggcctc gtgcaggctg aaaaacctga cacttctctg ctggccatgc tcctcagcgt 600
cctgctgcct ctgtcccccag gggccggcct ggccctgtgt tgctaccgac tcccaggagc 660
ccatctgcag cgtgcagctg ggggctgcag aagggaccct gcgtgcagtg gccccaaaga 720
tggccacacac agggaccacc cctggggcgg cgttccccc atggagtggg gccccacagc 780
cactggacag ccttggcccc tggaccctga gaactctcat gagccagcag gccaccctga 840
gaagcctctg ccagcagctc cgcctgaccc ccaagatcaa gtccagatgc caagatcctg 900
cctctgtgtg gaggttagct ctaaaatgaa cagatttaaa gacaggtggc cactgacagc 960
cactccagga ctaatgaactg cagggggcaga gccagtgaat caccggacct ccagcactct 1020
caggcagctt ggaagtcttc tcccagagt gagcttcgac ccaccacact caggaacca 1080
gagccacatt agaattctc gagggctgga gaaactgctc gggcacactc tccagctcaa 1140
taaaccatca gtcc 1154

```

<210> 362

<211> 953

<212> DNA

<213> Homo sapiens

<400> 362

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cagtggctac tgctgggatg gcgcatgtcc cacgctggag cagcagtgcc agcagctctg 60
ggggcctgat gggccaggaag tgacttctgc gggagccttg gcactcccac gtgccagct 120
ggacctgctt ggcctgggcc tggtagagcc aggcacccag tgtggacctt gaatggtgtg 180
ccagagcagg cgtgcaggga agaattgcct ccaggagctt cagcgtctgc tgactgcctg 240
ccacagccac ggggtttgca atagcaacca taactgccac cagcatggac agtggccctg tgcaggctga 300
accctctgtg gacaagccag gctttggtgg cagcatggac agtggccctg tgcaggctga 360
aaacctgac accctcctg tggccatgct cctcagcgtc ctgctgcctc tgctcccagg 420
ggccggcctg gctctgtgtt gctacogact cccaggagcc catctgcagc gatgcagctg 480
gggctgcaga agggaccctg cgtgcagttg ccccaaagat ggcccacaca gggaccaccc 540
cctggggcgg gttcacccca tggagttggg cccacagccc actggacagc cctggccctc 600
ggaccctcag aactctcatg agcccagcag ccaccctgag aagcctctgc cagcagcttc 660
gcctgacccc caagatcaag tccagatgcc aagatcctgc ctctggttag aggtagctcc 720
taaaatgaac agatttaaac acaggtggcc actgacagcc actccaggaa cttgaactgc 780
aggggcagag cagggaatc accggacctc cagcacctgc aggcagctt gaggtttctt 840
ccccgagtgg agcttcgacc caccactcc aggaacccag agccacatta gaagtctctg 900

```

agggctggag aacactgctg ggcacactct ccagctcaat aaaccatcag tcc

953

<210> 363

<211> 812

<212> PRT

<213> Homo sapiens

<400> 363

Met Gly Trp Arg Pro Arg Arg Ala Arg Gly Thr Pro Leu Leu Leu Leu
1 5 10 15

Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
20 25 30

Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
35 40 45

Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
50 55 60

Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
65 70 75 80

Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
85 90 95

Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp
100 105 110

His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val
115 120 125

Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg
130 135 140

Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp
145 150 155 160

Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
165 170 175

Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser
180 185 190

Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr
195 200 205

Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu
210 215 220

Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val
225 230 235 240

Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala
245 250 255

Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr
 260 265 270
 Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly
 275 280 285
 Leu Trp Ala Gln Arg Pro His Asp Ser Ala Gln Leu Leu Thr Gly Arg
 290 295 300
 Ala Phe Gln Gly Ala Thr Val Gly Leu Ala Pro Val Glu Gly Met Cys
 305 310 315 320
 Arg Ala Glu Ser Ser Gly Gly Val Ser Thr Asp His Ser Glu Leu Pro
 325 330 335
 Ile Gly Ala Ala Ala Thr Met Ala His Glu Ile Gly His Ser Leu Gly
 340 345 350
 Leu Ser His Asp Pro Asp Gly Cys Cys Val Glu Ala Ala Ala Glu Ser
 355 360 365
 Gly Gly Cys Val Met Ala Ala Ala Thr Gly His Pro Phe Pro Arg Val
 370 375 380
 Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Ala Phe Phe Arg Lys Gly
 385 390 395 400
 Gly Gly Ala Cys Leu Ser Asn Ala Pro Asp Pro Gly Leu Pro Val Pro
 405 410 415
 Pro Ala Leu Cys Gly Asn Gly Phe Val Glu Ala Gly Glu Glu Cys Asp
 420 425 430
 Cys Gly Pro Gly Gln Glu Cys Arg Asp Leu Cys Cys Phe Ala His Asn
 435 440 445
 Cys Ser Leu Arg Pro Gly Ala Gln Cys Ala His Gly Asp Cys Cys Val
 450 455 460
 Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu Cys Arg Gln Ala Met Gly
 465 470 475 480
 Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Ser His Cys Pro
 485 490 495
 Pro Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Arg Gly Ser Gly
 500 505 510
 Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys Gln Gln
 515 520 525
 Leu Trp Gly Pro Gly Ser His Pro Ala Pro Glu Ala Cys Phe Gln Val
 530 535 540
 Val Asn Ser Ala Gly Asp Ala His Gly Asn Cys Gly Gln Asp Ser Glu
 545 550 555 560

Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu
 565 570 575
 Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met Val Pro
 580 585 590
 Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys Arg Gly
 595 600 605
 Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu
 610 615 620
 Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg
 625 630 635 640
 Arg Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala
 645 650 655
 Cys His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala
 660 665 670
 Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser
 675 680 685
 Met Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu
 690 695 700
 Ala Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu
 705 710 715 720
 Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser
 725 730 735
 Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro
 740 745 750
 His Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro
 755 760 765
 Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu
 770 775 780
 Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro
 785 790 795 800
 Gln Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp
 805 810